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0425

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## RAW SEQUENCE LISTING

DATE: 04/24/2002

PATENT APPLICATION: US/10/036,492

TIME: 14:28:08

Input Set : A:\217943US0XCONT.txt

Output Set: N:\CRF3\04242002\J036492.raw

p. 6

ENTERED

3 <110> APPLICANT: HEMERLY, ADRIANA  
 4 FERREIRA, PAULO  
 5 ROMBAUTS, STEPHANE  
 7 <120> TITLE OF INVENTION: PLANT DNA REPLICATION MODULATING PROTEINS  
 9 <130> FILE REFERENCE: 217943US0XCONT  
 11 <140> CURRENT APPLICATION NUMBER: US 10/036,492  
 12 <141> CURRENT FILING DATE: 2002-01-07  
 14 <150> PRIOR APPLICATION NUMBER: EP 99202214.5  
 15 <151> PRIOR FILING DATE: 1999-07-05  
 17 <160> NUMBER OF SEQ ID NOS: 35  
 19 <170> SOFTWARE: PatentIn version 3.1  
 21 <210> SEQ ID NO: 1  
 22 <211> LENGTH: 889  
 23 <212> TYPE: PRT  
 24 <213> ORGANISM: Arabidopsis thaliana  
 26 <400> SEQUENCE: 1  
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 29 1 5 10 15  
 32 Arg Glu Leu Ile Pro Leu Ser Pro Thr Asn Ser Asp Gly Asn Asp Asp  
 33 20 25 30  
 36 Leu Asn Tyr His Leu His Ala Phe Glu Leu Ser Arg Leu Leu Leu Ser  
 37 35 40 45  
 40 Ser Gly His Pro Glu Ser Val Ile Asp Leu Ser Ser Lys Cys Thr Tyr  
 41 50 55 60  
 44 Phe Gln Gly Ser Pro Asn Leu Val Lys Tyr Leu Cys Ser Ile Pro Asn  
 45 65 70 75 80  
 48 Ser Pro Ile Ser Leu Ala Glu Asp Gly Phe Thr Val Thr Leu Ser Pro  
 49 85 90 95  
 52 Glu Ser Pro Ser Ala Pro Ala Ser Phe Ala Cys Ser Leu Asp Leu Gln  
 53 100 105 110  
 56 Glu Asn Val Val Leu Glu Gln Phe Met Asp Pro Arg Ser Leu Thr Leu  
 57 115 120 125  
 60 Lys His Ser Arg Glu Asn Ala Glu Gln Glu Glu Leu Glu Leu Met Pro  
 61 130 135 140  
 64 Leu Pro Lys Arg Ser Arg Asn Asp Gly Asn Asp Val Asn Tyr Ser Val  
 65 145 150 155 160  
 68 Ile Asp Ser Arg Pro Asn Asp Ile Arg Thr Val Ala Cys Gly Thr Met  
 69 165 170 175  
 72 Leu Gly Thr Ile Leu Ala Leu Glu Ser Gln Ala Ser Val Phe Asn Leu  
 73 180 185 190  
 76 Ser Ala Ser Asn Arg Gly Ile Glu Ala Phe Val Gln Asp His Gln Pro  
 77 195 200 205  
 80 Gly Pro Gln Thr Ser Asn Ala Ser Val Asp Val Asn Pro Thr His Arg

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81      210      215      220
84 Leu Glu Glu Ser Lys Asn Asp Leu Pro Ser Pro Gln Glu Asp Gly Tyr
85 225      230      235      240
88 Tyr Glu Arg Pro Glu Ile Gly Asp Phe Gln Ile Ala Asp Asn Gln Ile
89      245      250      255
92 Leu Ile Glu Glu Gly Asp Asp Lys Asn Lys Lys Asp Leu Phe Pro Lys
93      260      265      270
96 Gly Glu Ile Gln Thr Asp Ser Val Gln Ser Asp Pro Val Ala Ser Leu
97      275      280      285
100 Met Pro Thr Glu Asn Glu Leu Glu Pro Val Gln Ile Val Asp Asp Thr
101      290      295      300
104 Glu Asp Leu Leu Val Asp Asp His Thr Val Asp Ile Val Ser Thr Pro
105 305      310      315      320
108 Asp Arg Glu Leu Pro Leu Lys Pro Ser Ala Thr Glu Ala Asn Gln Asp
109      325      330      335
112 Lys Ser Leu Val Gln Lys Thr Leu Asp Gln Cys Lys Leu Pro Gly Asn
113      340      345      350
116 Ser Lys Thr Tyr Ser Cys Ser Pro Glu Ile Lys His Thr Arg Lys Ser
117      355      360      365
120 Lys Val Ile Gln Lys Arg Lys Gln Asn Phe Asn Thr Val Arg Leu Lys
121      370      375      380
124 Asp Gln Lys Asp Gln Ala Lys His Asn Thr Ile Pro Asp Phe Asp Ser
125 385      390      395      400
128 Tyr Thr Ile Val Glu Glu Glu Gly Ser Gly Gly Tyr Gly Ile Val Tyr
129      405      410      415
132 Lys Ala Thr Arg Lys Thr Asp Gly Thr Glu Phe Ala Ile Lys Cys Pro
133      420      425      430
136 His Val Gly Ala Gln Lys Tyr Tyr Val Asn Asn Glu Ile Arg Met Leu
137      435      440      445
140 Glu Arg Phe Gly Gly Lys Asn Cys Ile Ile Lys His Glu Gly Cys Leu
141      450      455      460
144 Lys Asn Gly Asp Ser Asp Cys Ile Ile Leu Glu His Leu Glu His Asp
145 465      470      475      480
148 Arg Pro Asp Ser Leu Lys Arg Glu Ile Asp Val Tyr Gln Leu Gln Trp
149      485      490      495
152 Tyr Gly Tyr Cys Met Phe Lys Ala Leu Ser Ser Leu His Lys Gln Gly
153      500      505      510
156 Val Val His Arg Asp Val Lys Pro Gly Asn Phe Leu Phe Ser Arg Lys
157      515      520      525
160 Thr Asn Lys Gly Tyr Leu Ile Asp Phe Asn Leu Ala Met Asp Leu His
161      530      535      540
164 Gln Lys Tyr Arg Arg Ala Asp Lys Ser Lys Ala Ala Ser Gly Leu Pro
165 545      550      555      560
168 Thr Ala Ser Lys Lys His His Thr Leu Val Lys Ser Leu Asp Ala Val
169      565      570      575
172 Asn Arg Gly Thr Asn Lys Pro Ser Gln Lys Thr Leu Ala Pro Asn Ser
173      580      585      590
176 Ile Lys Lys Ala Ala Gly Lys Thr Arg Ala Arg Asn Asp Met Thr Arg
177      595      600      605

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Input Set : A:\217943US0XCONT.txt

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180 Trp Glu Arg Leu Asn Ser Gln Gly Ala Glu Gly Ser Gly Leu Thr Ser
181      610                      615                      620
184 Ala Lys Asp Val Thr Ser Thr Arg Asn Asn Pro Ser Gly Glu Lys Arg
185 625                      630                      635                      640
188 Arg Glu Pro Leu Pro Cys His Gly Arg Lys Ala Leu Leu Asp Phe Leu
189                      645                      650                      655
192 Gln Glu Thr Met Ser Val Pro Ile Pro Asn His Glu Val Ser Ser Lys
193                      660                      665                      670
196 Ala Pro Thr Ser Met Arg Lys Arg Val Ala Ala Leu Pro Gly Lys Ala
197                      675                      680                      685
200 Glu Lys Glu Leu Leu Tyr Leu Thr Pro Met Pro Leu Cys Ser Asn Gly
201      690                      695                      700
204 Arg Pro Glu Ala Gly Asp Val Ile Glu Lys Lys Asp Gly Pro Cys Ser
205 705                      710                      715                      720
208 Gly Thr Lys Gly Phe Arg Ala Pro Glu Val Cys Phe Arg Ser Leu His
209                      725                      730                      735
212 Gln Gly Pro Lys Ile Asp Val Trp Ser Ala Gly Val Thr Leu Leu Tyr
213                      740                      745                      750
216 Leu Ile Met Gly Arg Thr Pro Phe Thr Gly Asp Pro Glu Gln Asn Ile
217                      755                      760                      765
220 Lys Asp Ile Ala Gln Leu Arg Gly Ser Glu Glu Leu Trp Glu Val Ala
221      770                      775                      780
224 Lys Leu His Asn Arg Glu Ser Ser Phe Pro Lys Glu Leu Tyr Glu Ser
225 785                      790                      795                      800
228 Arg Tyr Leu Lys Gly Met Glu Leu Arg Lys Trp Cys Glu Leu Asn Thr
229                      805                      810                      815
232 Lys Arg Arg Glu Phe Leu Asp Val Ile Pro Leu Ser Leu Leu Asp Leu
233                      820                      825                      830
236 Val Asp Lys Cys Leu Thr Val Asn Pro Arg Arg Arg Ile Ser Ala Glu
237      835                      840                      845
240 Asp Ala Leu Lys His Asp Phe His Pro Val His Glu Thr Leu Arg
241      850                      855                      860
244 Asn Gln Met Leu Leu Lys Gln Gln Pro Thr Val Val Ala Asp Ala Val
245 865                      870                      875                      880
248 Ser Gln Thr Leu Asn Tyr Leu Gln Leu
249                      885
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253 <211> LENGTH: 20
254 <212> TYPE: PRT
255 <213> ORGANISM: Artificial Sequence
257 <220> FEATURE:
258 <223> OTHER INFORMATION: Synthetic Peptide
260 <400> SEQUENCE: 2
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263 1                      5                      10                      15
266 Phe Ala Ile Lys
267      20
270 <210> SEQ ID NO: 3
271 <211> LENGTH: 20

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## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/036,492

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Input Set : A:\217943US0XCONT.txt

Output Set: N:\CRF3\04242002\J036492.raw

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272 <212> TYPE: PRT
273 <213> ORGANISM: Artificial Sequence
275 <220> FEATURE:
276 <223> OTHER INFORMATION: Synthetic Peptide
278 <400> SEQUENCE: 3
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281 1          5          10          15
284 Arg Ala Pro Glu
285          20
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289 <211> LENGTH: 29
290 <212> TYPE: PRT
291 <213> ORGANISM: Artificial Sequence
293 <220> FEATURE:
294 <223> OTHER INFORMATION: Synthetic Peptide
296 <400> SEQUENCE: 4
298 Asn Ile Lys Asp Ile Ala Gln Leu Arg Gly Ser Glu Glu Leu Trp Glu
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303          20          25
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307 <211> LENGTH: 728
308 <212> TYPE: PRT
309 <213> ORGANISM: Arabidopsis thaliana
311 <400> SEQUENCE: 5
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317 Phe Met Phe Thr Asn Ala Ile Phe Leu Cys Glu Leu Leu Leu Ala Gln
318          20          25          30
321 Phe Pro Ser Glu Val Asn Leu Gln Leu Leu Ala Arg Cys Tyr Leu Ser
322          35          40          45
325 Asn Ser Gln Ala Tyr Ser Ala Tyr Tyr Ile Leu Lys Gly Ser Lys Thr
326          50          55          60
329 Pro Gln Ser Arg Tyr Leu Phe Ala Phe Ser Cys Phe Lys Leu Asp Leu
330 65          70          75          80
333 Leu Gly Glu Ala Glu Ala Ala Leu Leu Pro Cys Glu Asp Tyr Ala Glu
334          85          90          95
337 Glu Val Pro Gly Gly Ala Ala Gly His Tyr Leu Leu Gly Leu Ile Tyr
338          100         105         110
341 Arg Tyr Ser Gly Arg Lys Asn Cys Ser Ile Gln Gln Phe Arg Met Ala
342          115         120         125
345 Leu Ser Phe Asp Pro Leu Cys Trp Glu Ala Tyr Gly Glu Leu Cys Ser
346          130         135         140
349 Leu Gly Ala Ala Glu Glu Ala Ser Thr Val Phe Gly Asn Val Ala Ser
350 145         150         155         160
353 Gln Arg Leu Gln Lys Thr Cys Val Glu Gln Arg Ile Ser Phe Ser Glu
354          165         170         175
357 Gly Ala Thr Ile Asp Gln Ile Thr Asp Ser Asp Lys Ala Leu Lys Asp
358          180         185         190

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Input Set : A:\217943US0XCONT.txt

Output Set: N:\CRF3\04242002\J036492.raw

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361 Thr Gly Leu Ser Gln Thr Glu His Ile Pro Gly Glu Asn Gln Gln Asp
362      195      200      205
365 Leu Lys Ile Met Gln Gln Pro Gly Asp Ile Pro Pro Asn Thr Asp Arg
366      210      215      220
369 Gln Leu Ser Thr Asn Gly Trp Asp Leu Asn Thr Pro Ser Pro Val Leu
370 225      230      235      240
373 Leu Gln Val Met Asp Ala Leu Pro Pro Leu Leu Leu Lys Asn Met Arg
374      245      250      255
377 Arg Pro Ala Val Glu Gly Ser Leu Met Ser Val His Gly Val Arg Val
378      260      265      270
381 Arg Arg Arg Asn Phe Phe Ser Glu Glu Leu Ser Ala Glu Ala Gln Glu
382      275      280      285
385 Glu Ser Gly Arg Arg Arg Ser Ala Arg Ile Ala Ala Arg Lys Lys Asn
386      290      295      300
389 Pro Met Ser Gln Ser Phe Gly Lys Asp Ser His Trp Leu His Leu Ser
390 305      310      315      320
393 Pro Ser Glu Ser Asn Tyr Ala Pro Ser Leu Ser Ser Met Ile Gly Lys
394      325      330      335
397 Cys Arg Ile Gln Ser Ser Lys Glu Val Ile Pro Asp Thr Val Thr Leu
398      340      345      350
401 Asn Asp Pro Ala Thr Thr Ser Gly Gln Ser Val Ser Asp Ile Gly Ser
402      355      360      365
405 Ser Val Asp Asp Glu Glu Lys Ser Asn Pro Ser Glu Ser Ser Pro Asp
406      370      375      380
409 Arg Phe Ser Leu Ile Ser Gly Ile Ser Glu Val Leu Ser Leu Leu Lys
410 385      390      395      400
413 Ile Leu Gly Asp Gly His Arg His Leu His Met Tyr Lys Cys Gln Glu
414      405      410      415
417 Ala Leu Leu Ala Tyr Gln Lys Leu Ser Gln Lys Gln Tyr Asn Thr His
418      420      425      430
421 Trp Val Leu Met Gln Val Gly Lys Ala Tyr Phe Glu Leu Gln Asp Tyr
422      435      440      445
425 Phe Asn Ala Asp Ser Ser Phe Thr Leu Ala His Gln Lys Tyr Pro Tyr
426      450      455      460
429 Ala Leu Glu Gly Met Asp Thr Tyr Ser Thr Val Leu Tyr His Leu Lys
430 465      470      475      480
433 Glu Glu Met Arg Leu Gly Tyr Leu Ala Gln Glu Leu Ile Ser Val Asp
434      485      490      495
437 Arg Leu Ser Pro Glu Ser Trp Cys Ala Val Gly Asn Cys Tyr Ser Leu
438      500      505      510
441 Arg Lys Asp His Asp Thr Ala Leu Lys Met Phe Gln Arg Ala Ile Gln
442      515      520      525
445 Leu Asn Glu Arg Phe Thr Tyr Ala His Thr Leu Cys Gly His Glu Phe
446      530      535      540
449 Ala Ala Leu Glu Glu Phe Glu Asp Ala Glu Arg Cys Tyr Arg Lys Ala
450 545      550      555      560
453 Leu Gly Ile Asp Thr Arg His Tyr Asn Ala Trp Tyr Gly Leu Gly Met
454      565      570      575
457 Thr Tyr Leu Arg Gln Glu Lys Phe Glu Phe Ala Gln His Gln Phe Gln

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RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/036,492

DATE: 04/24/2002  
TIME: 14:28:10

Input Set : A:\217943US0XCONT.txt  
Output Set: N:\CRF3\04242002\J036492.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:16; Xaa Pos. 1,2,3,5,6,9,10,12,13,14,15,16,17,18,19,21,22,23,25,26,28  
Seq#:16; Xaa Pos. 29,30,31,33,34  
Seq#:18; N Pos. 9,15,18  
Seq#:20; N Pos. 6,9,18  
Seq#:21; N Pos. 3,6,15  
Seq#:32; N Pos. 6,12  
Seq#:33; N Pos. 6,15  
Seq#:34; N Pos. 15  
Seq#:35; N Pos. 9